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Sequence Listing was accepted.

If you need help call the Patent Electronic Business Center at (866)
217-9197 (toll free).

Reviewer: Anne Corrigan

Timestamp: Sun Sep 09 17:25:59 EDT 2007

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Application No: 10822591 Version No: 2.0

Input Set:

Output Set:

Started: 2007-08-27 19:50:16.094
Finished: 2007-08-27 19:50:16.290
Elapsed: 0 hr(s) 0 min(s) 0 sec(s) 196 ms
Total Warnings: 3
Total Errors: 0
No. of SeqIDs Defined: 5
Actual SeqID Count: 5

Error code	Error Description
W 213	Artificial or Unknown found in <213> in SEQ ID (1)
W 213	Artificial or Unknown found in <213> in SEQ ID (2)
W 213	Artificial or Unknown found in <213> in SEQ ID (3)

SEQUENCE LISTING

<110> Leonard, Jack L.
Farwell, Alan P.

<120> NON-NUCLEAR EFFECTS OF THYROID HORMONE

<130> 07917-103002

<140> 10822591

<141> 2004-04-12

<150> US 09/894,734

<151> 2001-06-28

<150> US 60/214,572

<151> 2000-06-28

<160> 5

<170> FastSEQ for Windows Version 4.0

<210> 1

<211> 22

<212> PRT

<213> Artificial Sequence

<220>

<223> Synthetically generated peptide

<400> 1

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Gly	Phe	Ile	Ala	Arg	Val										
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<210> 2

<211> 18

<212> DNA

<213> Artificial Sequence

<220>

<223> Primer

<400> 2

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18

<210> 3

<211> 17

<212> DNA

<213> Artificial Sequence

<220>

<223> Primer

<400> 3

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17

<210> 4

<211> 1228

<212> DNA

<213> Homo sapiens

<400> 4

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gtcaagcggg agcagctcaa gaatggcggc ctgggcgtag tctccgacgc catctttgaa 180
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gcaggccaaa catggccaga ctcccttgct ttttgctgtg tagttccctc tgcctgggat 960
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gagtaggtac tccataaaag gtgtgttg 1228
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<210> 5

<211> 249

<212> PRT

<213> Homo sapiens

<400> 5

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          20          25          30
Leu Thr Leu Ser Gly Glu Met Ala Val Lys Arg Glu Gln Leu Lys Asn
          35          40          45
Gly Gly Leu Gly Val Val Ser Asp Ala Ile Phe Glu Leu Gly Lys Ser
          50          55          60
Leu Ser Ala Phe Asn Leu Asp Asp Thr Glu Val Ala Leu Leu Gln Ala
65          70          75          80
Val Leu Leu Met Ser Thr Asp Arg Ser Gly Leu Leu Cys Val Asp Lys
          85          90          95
Ile Glu Lys Ser Gln Glu Ala Tyr Leu Leu Ala Phe Glu His Tyr Val
          100          105          110
Asn His Arg Lys His Asn Ile Pro His Phe Trp Pro Lys Leu Leu Met
          115          120          125
Lys Glu Arg Glu Val Gln Ser Ser Ile Leu Tyr Lys Gly Ala Ala Ala
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130		135		140	
Glu Gly Arg Pro Gly Gly Ser Leu Gly Val His Pro Glu Gly Gln Gln					
145		150		155	160
Leu Leu Gly Met His Val Val Gln Gly Pro Gln Val Arg Gln Leu Glu					
	165		170		175
Gln Gln Leu Gly Glu Ala Gly Ser Leu Gln Gly Pro Val Leu Gln His					
	180		185		190
Gln Ser Pro Lys Ser Pro Gln Gln Arg Leu Leu Glu Leu Leu His Arg					
	195		200		205
Ser Gly Ile Leu His Ala Arg Ala Val Cys Gly Glu Asp Asp Ser Ser					
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Glu Ala Asp Ser Pro Ser Ser Ser Glu Glu Glu Pro Glu Val Cys Glu					
225		230		235	240
Asp Leu Ala Gly Asn Ala Ala Ser Pro					
	245				